

4 Thu May 2 11:45:17 2002

us-09-770-693-3.rapm

1brahim M.  
09/7706933 Page 1  
Seq. IDs 3 of 4 w/  
Interf

GenScope version 4.5  
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OM protein - protein search, using sw model

Run on: May 2, 2002, 03:27:49 | Search time 126.22 seconds  
1095.460 Million cell updates/sec

Title: US-09-770-693-3  
Perfect score: 2079  
Sequence: 1 NLSFSGISGASTMSTIGGA.....DAMDAKLNINLALZAA 403

Scoring table:

BLOSUM62

Gap: 10.0, Gapext 0.5

Search: 3516493 seqs, 35124056 residues

Total number of hits satisfying chosen parameters: 3516493

Minimum DB seq length: 0

Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 43 summaries

Database:

- 1: /cni2\_6/prodata/2/paa/US88\_COMB.pep.\*
- 2: /cni2\_6/prodata/2/paa/US88\_COMB.pep.\*
- 3: /cni2\_6/prodata/2/paa/US88\_COMB.pep.\*
- 4: /cni2\_6/prodata/2/paa/US88\_COMB.pep.\*
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- 32: /cni2\_6/prodata/2/paa/US88\_COMB.pep.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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1	2079	100.0	403	1 PCT-0501-02579-3 Sequence 3, Appl

2	2079	100.0	403	1 PCT-0597-22629-3 Sequence 3, Appl
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4	2079	100.0	403	1 PCT-0598-01507-3 Sequence 3, Appl
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Matches 403: Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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 Db 1 NSLWFGSGASTWQISIGAGGNNGLGTSRQMGGLGNSALGSGNMGWYVWKLGL 60

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Qy 121 TTTSTSTPDLALGISTNSQMSDSTSTSTSTSTSTSTSTSTSTSTSTSTSTST 180  
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 Db 241 GKGGLGSGVDTGGLGAGVGTGKAGGLGGLGGLGGLGGLGGLGGLGGLGGLGGL 300

Qy 301 GQPMQGPVFPVGRVYKQVGTGKSWALSKFQDQDGFASDFYKMGDKMKEI 360  
 Db 301 GQPMQGPVFPVGRVYKQVGTGKSWALSKFQDQDGFASDFYKMGDKMKEI 360

Qy 361 PMAGDTGNGGAGGSSGLGDMMAGGATNNALGGLGAA 403  
 Db 361 PMAGDTGNGGAGGSSGLGDMMAGGATNNALGGLGAA 403

RESULT 2  
 : Sequence 3, Application PC/TUS9722629  
 : GENERAL INFORMATION: Research Foundation, Inc.  
 : TITLE OF INVENTION: HYPERSENSITIVE RESPONSE INDICED  
 : TITLE OF INVENTION: RESISTANCE IN PLANTS BY SEED TREATMENT  
 : INVENTOR: Goldan, Michael L.  
 : CORRESPONDENCE ADDRESS:  
 : ADDRESSEE: Nixon, Hargrave, Devans & Doyle LLP  
 : ATTORNEY/AGENT INFORMATION:  
 : CITY: Rochester  
 : STATE: New York  
 : ZIP: 14603  
 : COMPUTER READABLE FORM:  
 : OPERATING SYSTEM: IBM PC compatible  
 : CURRENT APPLICATION IN RELEASE #1.0, Version #1.30  
 : APPLICATION NUMBER: PCT/US97/22629  
 : FILING DATE:  
 : PRIOR APPLICATION DATA:  
 : APPLICATION NUMBER: US 60/033,230  
 : INVENTOR/AGENT INFORMATION:  
 : NAME: Goldan, Michael L.  
 : REFERENCE/DOCKET NUMBER: 12603/1202  
 : TELECOMMUNICATION INFORMATION:  
 : TELEPHONE: (716) 263-1004  
 : TELEFAX: (716) 263-1004  
 : INFORMATION FOR SEQ ID NO: 3:  
 : SEQUENCE CHARACTERISTICS:  
 : TYPE: amino acid  
 : STRANDNESS:  
 : MOLECULE TYPE: Protein  
 : PCT-US97-22629-3

Query Match 100.0% Score 2079; Db 1; Length 403;  
 Best Local Similarity 100.0% Ident. No. 1; de-169;  
 Matches 403: Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy 121 TTTSTSTPDLALGISTNSQMSDSTSTSTSTSTSTSTSTSTSTSTSTSTSTST 180  
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Qy 361 PMAGDTGNGGAGGSSGLGDMMAGGATNNALGGLGAA 403  
 Db 361 PMAGDTGNGGAGGSSGLGDMMAGGATNNALGGLGAA 403

RESULT 3  
 : Sequence 3, Application PC/TUS9801507  
 : GENERAL INFORMATION: Cornell Research Foundation, Inc.  
 : TITLE OF INVENTION: ENHANCEMENT OF GROWTH IN PLANTS  
 : INVENTOR: Goldan, Michael L.  
 : CORRESPONDENCE ADDRESS:  
 : ADDRESSEE: Nixon, Hargrave, Devans & Doyle LLP  
 : ATTORNEY/AGENT INFORMATION:  
 : CITY: Rochester  
 : STATE: New York  
 : ZIP: 14603  
 : COMPUTER READABLE FORM:  
 : OPERATING SYSTEM: IBM PC compatible  
 : CURRENT APPLICATION IN RELEASE #1.0, Version #1.30  
 : APPLICATION NUMBER: PCT/US98/01507  
 : FILING DATE:  
 : PRIOR APPLICATION DATA:  
 : APPLICATION NUMBER: US 60/096,048  
 : INVENTOR/AGENT INFORMATION:  
 : NAME: Goldan, Michael L.  
 : REFERENCE/DOCKET NUMBER: 19603/1502  
 : TELECOMMUNICATION INFORMATION:  
 : TELEPHONE: (716) 263-1004  
 : TELEFAX: (716) 263-1004  
 : INFORMATION FOR SEQ ID NO: 3:  
 : SEQUENCE CHARACTERISTICS:  
 : TYPE: amino acid  
 : STRANDNESS:  
 : MOLECULE TYPE: Protein

TOPLOGY: linear  
MOLECULE TYPE: protein  
PCT-US98-01507-3

Query Match 100.0%; Score 2079; DB 1; Length 403;  
Best Local Similarity 100.0%; Pred. No. 1,4e+169; Gaps 0;  
Matches 403; Conservative 0; Indels 0;  
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DB 1 MSMTSGLSGNTQWQIGTGCAGGNGNLLGTSRQNGLSGSLGAGGQNDQVYGLAGLL 60  
QY 61 TMMHMMHMMGGGGLAGGGLAGGGLAGGGLAGGGLAGGGLAGGGLAGGGLAGGGLAGG 120  
DB 61 TMMHMMHMMGGGGLAGGGLAGGGLAGGGLAGGGLAGGGLAGGGLAGGGLAGGGLAGG 120  
QY 121 TSTTNSPLDQALGINTSQNDQSTGTSSTSSSDPQQQLAMFSEHMSIFGQDQDT 180  
DB 121 TSTTNSPLDQALGINTSQNDQSTGTSSTSSSDPQQQLAMFSEHMSIFGQDQDT 180  
QY 181 QSSSGSGGQPTREQDQVNTKRYTALSGNLGSLGSLGAGGGLAGGGLAGGGLAGG 240  
DB 181 QSSSGSGGQPTREQDQVNTKRYTALSGNLGSLGSLGAGGGLAGGGLAGGGLAGG 240  
QY 241 GGGELQNLGSGDQVQLANAVGTQGMAGTQALMDGTHRSSTFRFYKNGDRAKAEI 300  
DB 241 GGGELQNLGSGDQVQLANAVGTQGMAGTQALMDGTHRSSTFRFYKNGDRAKAEI 300  
QY 301 GQPMQYFVETKPYQKPGQGVETDQSKALKSPDDQDQMTASHEQFNKAGHMR 360  
DB 301 GQPMQYFVETKPYQKPGQGVETDQSKALKSPDDQDQMTASHEQFNKAGHMR 360  
QY 361 PHAGDTGNGNLQNGAGGSSSLGTDMMAGDIAINNALGKLGAA 403  
DB 361 PHAGDTGNGNLQNGAGGSSSLGTDMMAGDIAINNALGKLGAA 403

RESULT 4  
PCT-US98-03604-3  
Sequence Information: Application PC/US9803604  
APPLICANT: Corneil Research Foundation, Inc.  
TITLE OF INVENTION: INSECT CONTROL WITH A.  
NUMBER OF SEQUENCES: 10  
CORRESPONDENCE ADDRESS: Debra A. DeGrave, Devoes & Doyle LLP  
STREET: P.O. Box 1051, Clinton Square  
CITY: Rochester  
STATE: New York  
ZIP: 14603  
COMPUTER READABLE FORM: disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/NE-DOS  
CURRENT APPLICATION DATA: US 80/439,226  
APPLICATION NUMBER: PCT/US98/03604  
FILING DATE: 08/09/98  
PRIORITY INFORMATION: US 80/439,226  
ANTONY/AGNT INFORMATION:  
NAME: Goldman, Michael L.  
REFERENCE/DOCUM NUMBER: 19603/4522  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (716) 263-1304  
FAX: (716) 263-1304  
INFORMATION FOR SEQ. ID NO: 3:

SEQUENCE CHARACTERISTICS:  
LENGTH: 403 amino acids  
TOPLOGY: linear  
MOLECULE TYPE: protein  
PCT-US98-03604-3

Query Match 100.0%; Score 2079; DB 1; Length 403;  
Best Local Similarity 100.0%; Pred. No. 1,4e+169; Gaps 0;  
Matches 403; Conservative 0; Indels 0;  
QY 1 MSMTSGLSGNTQWQIGTGCAGGNGNLLGTSRQNGLSGSLGAGGQNDQVYGLAGLL 60  
DB 1 MSMTSGLSGNTQWQIGTGCAGGNGNLLGTSRQNGLSGSLGAGGQNDQVYGLAGLL 60  
QY 61 TMMHMMHMMGGGGLAGGGLAGGGLAGGGLAGGGLAGGGLAGGGLAGGGLAGGGLAGG 120  
DB 61 TMMHMMHMMGGGGLAGGGLAGGGLAGGGLAGGGLAGGGLAGGGLAGGGLAGGGLAGG 120  
QY 121 TSTTNSPLDQALGINTSQNDQSTGTSSTSSSDPQQQLAMFSEHMSIFGQDQDT 180  
DB 121 TSTTNSPLDQALGINTSQNDQSTGTSSTSSSDPQQQLAMFSEHMSIFGQDQDT 180  
QY 181 QSSSGSGGQPTREQDQVNTKRYTALSGNLGSLGSLGAGGGLAGGGLAGGGLAGG 240  
DB 181 QSSSGSGGQPTREQDQVNTKRYTALSGNLGSLGSLGAGGGLAGGGLAGGGLAGG 240  
QY 241 GGGELQNLGSGDQVQLANAVGTQGMAGTQALMDGTHRSSTFRFYKNGDRAKAEI 300  
DB 241 GGGELQNLGSGDQVQLANAVGTQGMAGTQALMDGTHRSSTFRFYKNGDRAKAEI 300  
QY 301 GQPMQYFVETKPYQKPGQGVETDQSKALKSPDDQDQMTASHEQFNKAGHMR 360  
DB 301 GQPMQYFVETKPYQKPGQGVETDQSKALKSPDDQDQMTASHEQFNKAGHMR 360  
QY 361 PHAGDTGNGNLQNGAGGSSSLGTDMMAGDIAINNALGKLGAA 403  
DB 361 PHAGDTGNGNLQNGAGGSSSLGTDMMAGDIAINNALGKLGAA 403

RESULT 5  
PCT-US98-0376-2  
Sequence Information: Application US/08851376  
APPLICANT: Debra A. DeGrave, Devoes & Doyle LLP  
APPLICANT: Bauer, David W.  
APPLICANT: Debra A. DeGrave, Devoes & Doyle LLP  
APPLICANT: Debra A. DeGrave, Devoes & Doyle LLP  
APPLICANT: Debra A. DeGrave, Devoes & Doyle LLP  
TITLE OF INVENTION: ELICITOR OF THE HYPERSENSITIVE RESPONSE  
NUMBER OF SEQUENCES: 5  
CORRESPONDENCE ADDRESS: Debra A. DeGrave, Devoes & Doyle LLP  
STREET: Clinton Square, P.O. Box 1051  
CITY: Rochester  
STATE: New York  
ZIP: 14603  
COMPUTER READABLE FORM: disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/NE-DOS  
CURRENT APPLICATION DATA: US 80/439,226  
APPLICATION NUMBER: US/08/451,376  
FILING DATE: 08/09/98  
PRIORITY INFORMATION: US 80/439,226  
INFORMATION FOR SEQ. ID NO: 4:

? FILING DATE: 23-FEB-1994  
 ? ATTORNEY/AGENT INFORMATION:  
 ? NAME: Goldmar, Michael L.  
 ? REGISTRATION NUMBER: 30,727  
 ? REFERENCE/DOCKET NUMBER: 19603/10035  
 ? TELEPHONE: (716) 263-1304  
 ? TELEFAX: (716) 263-1600  
 ? INFORMATION FOR SEQ ID NO: 2:  
 ? LENGTH: 403 amino acids  
 ? TYPE: amino acid  
 ? TOPOLOGY: linear  
 ? MOLECULE TYPE: peptide  
 ? US-09-091-376-2

Query Match 100.0%; Score 2079; DB 12; Length 403;  
 Best Local Similarity 100.0%; Pred. No. 1; 66-169; Indels 0; Gaps 0;  
 Matches 403; Conservative 0; Mismatches 0;

QY 1 MSLSFSGTASWISGIGAGGNGGLGTSRQNGALGCGNGWVQAGLL 60  
 DB 1 MSLSFSGTASWISGIGAGGNGGLGTSRQNGALGCGNGWVQAGLL 60  
 QY 61 TQMMNMHMGWGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGL 120  
 DB 61 TQMMNMHMGWGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGL 120  
 QY 121 TTSSTNSPLDQALNTSRQDSTSTSDSPQMLLKFSEINGSLFGDQDQT 180  
 DB 121 TTSSTNSPLDQALNTSRQDSTSTSDSPQMLLKFSEINGSLFGDQDQT 180  
 QY 181 QSSSSGQPTEDQNYKGYDALSIMNGLSQLLNGGLGGLGGLGGLGGL 240  
 DB 181 QSSSSGQPTEDQNYKGYDALSIMNGLSQLLNGGLGGLGGLGGLGGL 240  
 QY 241 GQGLQNLSPDYVQYQJGNAVGTGIMKAGLQALNDGTHRSSTFSYNGDRAMKEI 300  
 DB 241 GQGLQNLSPDYVQYQJGNAVGTGIMKAGLQALNDGTHRSSTFSYNGDRAMKEI 300  
 QY 301 GQPMQYFVFPKQYQJGNAVGTGIMKAGLQALNDGTHRSSTFSYNGDRAMKEI 360  
 DB 301 GQPMQYFVFPKQYQJGNAVGTGIMKAGLQALNDGTHRSSTFSYNGDRAMKEI 360  
 QY 361 PMAGDTGNCNLARGAGGSSGGLTDANMAGDAINNALGEGAA 403  
 DB 361 PMAGDTGNCNLARGAGGSSGGLTDANMAGDAINNALGEGAA 403

? RESULT 6  
 ? US-09-030-270-3  
 ? Sequence 3, Application US/09030270  
 ? GENERAL INFORMATION:  
 ? APPLICANT: Zitter, Thomas A.  
 ? APPLICANT: Wei, Zhong-Min  
 ? TITLE OF INVENTION: HYPERSENSITIVE RESPONSE ELICITOR  
 ? NUMBER OF SEQUENCES: 10  
 ? COMPLETENCE: ADDRESS:  
 ? ADDRESS: Nixon, Hargrave, Devans & Doyle LLP  
 ? STREET: P.O. Box 1051, Clinton Square  
 ? CITY: Rochester  
 ? STATE: New York  
 ? COUNTRY: U.S.A.  
 ? ZIP: 14603  
 ? COMEDIN TYPE: floppy disk  
 ? COMPUTER: IBM PC compatible  
 ? SOFTWARE: Patent In Relates #1.0, Version #1.30  
 ? CURRENT APPLICATION DATA:

? APPLICATION NUMBER: US/09/030,270  
 ? FILING DATE:  
 ? PRIOR APPLICATION DATA:  
 ? APPLICATION NUMBER: US 6/039,226  
 ? FILING DATE: FEB-1997  
 ? ATTORNEY/AGENT INFORMATION:  
 ? NAME: Goldmar, Michael L.  
 ? REGISTRATION NUMBER: 30,727  
 ? REFERENCE/DOCKET NUMBER: 19603/10035  
 ? TELEPHONE: (716) 263-1304  
 ? TELEFAX: (716) 263-1600  
 ? INFORMATION FOR SEQ ID NO: 3:  
 ? LENGTH: 403 amino acids  
 ? TYPE: amino acid  
 ? TOPOLOGY: linear  
 ? MOLECULE TYPE: protein  
 ? US-09-030-270-3

Query Match 100.0%; Score 2084; DB 14; Length 403;  
 Best Local Similarity 100.0%; Pred. No. 1; 66-169; Indels 0; Gaps 0;  
 Matches 403; Conservative 0; Mismatches 0;

QY 1 MSLSFSGTASWISGIGAGGNGGLGTSRQNGALGCGNGWVQAGLL 60  
 DB 1 MSLSFSGTASWISGIGAGGNGGLGTSRQNGALGCGNGWVQAGLL 60  
 QY 61 TQMMNMHMGWGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGL 120  
 DB 61 TQMMNMHMGWGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGL 120  
 QY 121 TTSSTNSPLDQALNTSRQDSTSTSDSPQMLLKFSEINGSLFGDQDQT 180  
 DB 121 TTSSTNSPLDQALNTSRQDSTSTSDSPQMLLKFSEINGSLFGDQDQT 180  
 QY 181 QSSSSGQPTEDQNYKGYDALSIMNGLSQLLNGGLGGLGGLGGLGGL 240  
 DB 181 QSSSSGQPTEDQNYKGYDALSIMNGLSQLLNGGLGGLGGLGGLGGL 240  
 QY 241 GQGLQNLSPDYVQYQJGNAVGTGIMKAGLQALNDGTHRSSTFSYNGDRAMKEI 300  
 DB 241 GQGLQNLSPDYVQYQJGNAVGTGIMKAGLQALNDGTHRSSTFSYNGDRAMKEI 300  
 QY 301 GQPMQYFVFPKQYQJGNAVGTGIMKAGLQALNDGTHRSSTFSYNGDRAMKEI 360  
 DB 301 GQPMQYFVFPKQYQJGNAVGTGIMKAGLQALNDGTHRSSTFSYNGDRAMKEI 360  
 QY 361 PMAGDTGNCNLARGAGGSSGGLTDANMAGDAINNALGEGAA 403  
 DB 361 PMAGDTGNCNLARGAGGSSGGLTDANMAGDAINNALGEGAA 403

? RESULT 7  
 ? US-09-030-270-3  
 ? Sequence 23, Application US/09046118  
 ? GENERAL INFORMATION:  
 ? APPLICANT: Zitter, Ronald J.  
 ? APPLICANT: Wei, Zhong-Min  
 ? TITLE OF INVENTION: HYPERSENSITIVE RESPONSE ELICITOR  
 ? NUMBER OF SEQUENCES: 30  
 ? COMPLETENCE: ADDRESS:  
 ? ADDRESS: Nixon, Hargrave, Devans & Doyle LLP  
 ? STREET: Clinton Square, P.O. Box 1051  
 ? CITY: Rochester  
 ? STATE: New York  
 ? COUNTRY: U.S.A.

```

1 ZIP: 1603
2 COMPUTER: IBM PC compatible
3 MEDIA TYPE: Floppy disk
4 COMPUTER: IBM PC compatible
5 SOFTWARE: Patentin Release #1.0, Version #1.30
6 CURRENT APPLICATION DATA:
7 FILING DATE: 05/09/086.118
8 APPLICATION NUMBER: US/09/086.118
9 CLASSIFICATION:
10 PRIOR APPLICATION DATA: US 60/048,109
11 FILING DATE: 30-May-1997
12 ATTORNEY/AGENT INFORMATION:
13 REGISTRATION NUMBER: 30,727
14 REFERENCE/DOCKET NUMBER: 19603/1301
15 TELECOMMUNICATIONS INFORMATION:
16 TELEFAX: (716) 263-1600
17 INFORMATION FOR SEQ ID NO: 23:
18 SOURCE: amino acids
19 LENGTH: 403 amino acids
20 TYPE: amino acid
21 TOPOLOGY: linear
22 MOLECULE TYPE: protein
23 US-09-086-118-23

Query Match 100.0% Score 2079; db 18; Length 403;
Best Local Similarity 100.0%; Pred. No. 1,4e+169;
Matches 403; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MSLSLGLSGLSTWQISTGAGNGLLGTFRQNLGAGLGGNGLGCGNQNTVNLGLL 60
Db 1 MSLSLGLSGLSTWQISTGAGNGLLGTFRQNLGAGLGGNGLGCGNQNTVNLGLL 60
QY 61 TQMMHNSMNGGGLMGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGL 120
Db 61 TQMMHNSMNGGGLMGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGL 120
QY 121 TSTTNSFLDQALNSTSQNDISTSTQTSUSSDPMQLKMFSEIAGSLGKGN 180
Db 121 TSTTNSFLDQALNSTSQNDISTSTQTSUSSDPMQLKMFSEIAGSLGKGN 180
QY 122 TSTTNSFLDQALNSTSQNDISTSTQTSUSSDPMQLKMFSEIAGSLGKGN 180
Db 122 TSTTNSFLDQALNSTSQNDISTSTQTSUSSDPMQLKMFSEIAGSLGKGN 180
QY 181 GSSSSGGQPTQEDQATKAYTQALGKGLGGLGGLGGLGGLGGLGGLGGLGGL 240
Db 181 GSSSSGGQPTQEDQATKAYTQALGKGLGGLGGLGGLGGLGGLGGLGGLGGL 240
QY 241 GGGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGL 300
Db 241 GGGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGL 300
QY 301 GQPMQDPTVPTQYQKQPGQVDTQKSNALKSPDDQDPTFASHQDPAKAGMR 360
Db 301 GQPMQDPTVPTQYQKQPGQVDTQKSNALKSPDDQDPTFASHQDPAKAGMR 360
QY 361 PHAGDTGNGLGARGAGSSGLGDAMMAGDANNALGKGLAA 403
Db 361 PHAGDTGNGLGARGAGSSGLGDAMMAGDANNALGKGLAA 403

RESULT 0
US-09-431-614-3
: Sequence 3, Application US/09431614
: GENERAL INFORMATION:
: APPLICANT: Schindig, Richard L.
: TITLE OF INVENTION: HYPERSENSITIVE RESPONSE ELICITOR-INDUCED STRESS
: FILE REFERENCE: 21829/41 (SEC-003)
: CURRENT APPLICATION NUMBER: US/09/431.614
: EARLIER FILING DATE: 1998-11-05
: DARTER APPLICATION NUMBER: 60/107,243
: NUMBER OF SEQ ID NOS: 18
: SOURCE: Patentin Ver. 2.0
: SEQ ID NO 3
: LENGTH: 403
: MOLECULE TYPE: protein
: ORGANISM: Brinia amylovora
: US-09-431-614-3

Query Match 100.0% Score 2079; db 18; Length 403;
Best Local Similarity 100.0%; Pred. No. 1,4e+169;
Matches 403; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MSLSLGLSGLSTWQISTGAGNGLLGTFRQNLGAGLGGNGLGCGNQNTVNLGLL 60

```



Db 121 TSTTNSFLQALQKYNSTQNDISTQDTSQSSDPQQLKMFSEMQSLQDQDQT 180  
 QY 181 QSSSSGGQPTQEDQNAKYKQYDALSJGKGLQKGLQGLQGLQGLQGLQGLQGL 240  
 Db 181 QSSSSGGQPTQEDQNAKYKQYDALSJGKGLQKGLQGLQGLQGLQGLQGLQGL 240  
 QY 241 GSGGLQNSGPTQQLANAVGTQKQKQALQALQDGTTHBSSTSFYKCDRAKKEI 300  
 Db 241 GSGGLQNSGPTQQLANAVGTQKQKQALQALQDGTTHBSSTSFYKCDRAKKEI 300  
 QY 301 QDQMDQYFVFGPTQKQKQYQVTDQKMAKALSKFPDQDQMTFMSQFNAKGIKR 360  
 Db 301 QDQMDQYFVFGPTQKQKQYQVTDQKMAKALSKFPDQDQMTFMSQFNAKGIKR 360  
 QY 361 PMAGDTQNGNLQKQAGSSSGGTDNNMAGDAIINNMLKGLQAA 403  
 Db 361 PMAGDTQNGNLQKQAGSSSGGTDNNMAGDAIINNMLKGLQAA 403

RESULT 12  
 US-09-770-693-3  
 : Application US/09770693  
 : GENERAL INFORMATION:  
 : APPLICANT: Beer, Steven V.  
 : TITLE OF INVENTION: OXYCETE-RESISTANT TRANSGENIC PLANTS BY VIRTUE OF  
 : FILE OF INVENTION: PATOGEN-INDUCED EXPRESSION OF A RETROLOGOUS  
 : FILE REFERENCE: 1950/2501  
 : CURRENT APPLICATION NUMBER: US/09/770.693  
 : PRIORITY DATE: 2000-01-26  
 : PRIORITY FILING DATE: 2000-01-26  
 : NUMBER OF SEQ ID NOS: 26  
 : SOFTWARE: PATGEN Ver. 2.1  
 : SEQ ID NO 3  
 : LENGTH: 403  
 : ORGANISM: Erwinia amylovora  
 : US-09-770-693-3

Query Match 100.0% Score 2079; Db 21; Length 403;  
 Similarity 100.0% Percent of Identical Regions 66.66%  
 Matches 403; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MSLSNLSGASTQVSTSGAGQNGGLGATSSQNLGAGSLQGLQGLQGLQGLQGLQGL 60  
 Db 1 MSLSNLSGASTQVSTSGAGQNGGLGATSSQNLGAGSLQGLQGLQGLQGLQGLQGL 60  
 QY 61 TOMHHNHSNGSGGLQWGLQGLQGLQGLQGLQGLQGLQGLQGLQGLQGLQGLQGLN 120  
 Db 61 TOMHHNHSNGSGGLQWGLQGLQGLQGLQGLQGLQGLQGLQGLQGLQGLQGLN 120  
 QY 121 TSTTNSFLQALQKYNSTQNDISTQDTSQSSDPQQLKMFSEMQSLQDQDQT 180  
 Db 121 TSTTNSFLQALQKYNSTQNDISTQDTSQSSDPQQLKMFSEMQSLQDQDQT 180  
 QY 181 QSSSSGGQPTQEDQNAKYKQYDALSJGKGLQKGLQGLQGLQGLQGLQGLQGL 240  
 Db 181 QSSSSGGQPTQEDQNAKYKQYDALSJGKGLQKGLQGLQGLQGLQGLQGLQGL 240  
 QY 241 GSGGLQNSGPTQQLANAVGTQKQKQALQALQDGTTHBSSTSFYKCDRAKKEI 300  
 Db 241 GSGGLQNSGPTQQLANAVGTQKQKQALQALQDGTTHBSSTSFYKCDRAKKEI 300  
 QY 301 QDQMDQYFVFGPTQKQKQYQVTDQKMAKALSKFPDQDQMTFMSQFNAKGIKR 360  
 Db 301 QDQMDQYFVFGPTQKQKQYQVTDQKMAKALSKFPDQDQMTFMSQFNAKGIKR 360  
 QY 361 PMAGDTQNGNLQKQAGSSSGGTDNNMAGDAIINNMLKGLQAA 403  
 Db 361 PMAGDTQNGNLQKQAGSSSGGTDNNMAGDAIINNMLKGLQAA 403

RESULT 13  
 US-09-835-684-3  
 : Application US/09835684  
 : GENERAL INFORMATION:  
 : APPLICANT: Wei, Zhong-Min  
 : TITLE OF INVENTION: TREATMENT OF FRUITS OR VEGETABLES WITH HYPERSENSITIVE  
 : FILE OF INVENTION: RESPONSE ELICITOR TO CONTROL POSTHARVEST DISEASE OR  
 : FILE REFERENCE: 2182/71  
 : CURRENT APPLICATION NUMBER: US/09/835.684  
 : PRIORITY DATE: 2000-04-19  
 : PRIORITY FILING DATE: 2000-04-19  
 : NUMBER OF SEQ ID NOS: 26  
 : SOFTWARE: PATGEN Ver. 2.1  
 : SEQ ID NO 3  
 : LENGTH: 403  
 : ORGANISM: Erwinia amylovora  
 : US-09-835-684-3

Query Match 100.0% Score 2079; Db 22; Length 403;  
 Similarity 100.0% Percent of Identical Regions 66.66%  
 Matches 403; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MSLSNLSGASTQVSTSGAGQNGGLGATSSQNLGAGSLQGLQGLQGLQGLQGLQGL 60  
 Db 1 MSLSNLSGASTQVSTSGAGQNGGLGATSSQNLGAGSLQGLQGLQGLQGLQGLQGL 60  
 QY 61 TOMHHNHSNGSGGLQWGLQGLQGLQGLQGLQGLQGLQGLQGLQGLQGLQGLN 120  
 Db 61 TOMHHNHSNGSGGLQWGLQGLQGLQGLQGLQGLQGLQGLQGLQGLQGLQGLN 120  
 QY 121 TSTTNSFLQALQKYNSTQNDISTQDTSQSSDPQQLKMFSEMQSLQDQDQT 180  
 Db 121 TSTTNSFLQALQKYNSTQNDISTQDTSQSSDPQQLKMFSEMQSLQDQDQT 180  
 QY 181 QSSSSGGQPTQEDQNAKYKQYDALSJGKGLQKGLQGLQGLQGLQGLQGLQGL 240  
 Db 181 QSSSSGGQPTQEDQNAKYKQYDALSJGKGLQKGLQGLQGLQGLQGLQGLQGL 240  
 QY 241 GSGGLQNSGPTQQLANAVGTQKQKQALQALQDGTTHBSSTSFYKCDRAKKEI 300  
 Db 241 GSGGLQNSGPTQQLANAVGTQKQKQALQALQDGTTHBSSTSFYKCDRAKKEI 300  
 QY 301 QDQMDQYFVFGPTQKQKQYQVTDQKMAKALSKFPDQDQMTFMSQFNAKGIKR 360  
 Db 301 QDQMDQYFVFGPTQKQKQYQVTDQKMAKALSKFPDQDQMTFMSQFNAKGIKR 360  
 QY 361 PMAGDTQNGNLQKQAGSSSGGTDNNMAGDAIINNMLKGLQAA 403  
 Db 361 PMAGDTQNGNLQKQAGSSSGGTDNNMAGDAIINNMLKGLQAA 403

RESULT 14  
 US-09-879-248-3  
 : Application US/09879248  
 : GENERAL INFORMATION:  
 : APPLICANT: Wei, Zhong-Min  
 : TITLE OF INVENTION: HYPERSENSITIVE RESPONSE ELICITING DOMAINS AND USE  
 : FILE OF INVENTION: THEREOF  
 : FILE REFERENCE: 2182/81  
 : CURRENT APPLICATION NUMBER: US/09/879.248  
 : PRIORITY DATE: 2000-06-16  
 : PRIORITY FILING DATE: 2000-06-16  
 : NUMBER OF SEQ ID NOS: 18

! SOFTWARE: PatentIn Ver. 2.1  
! SEQ ID NO 3  
! TYPE: PRT  
! ORGANISM: Erwinia amylovora  
! US-09-679-248-3

Query Match 100.0% Score 2079; Db 22; Length 403;  
Best Local Similarity 100.0%; Pos 21; Indels 0; Gaps 0;  
Matches 403; Conservative 0; Mismatches 0;

Oy 1 NSLWTSGLASTPMQISIGAGNNGLLGTSRQNGALGAGGNGNTVWLAGLL 60  
Oy 61 TMMHMSGSGGGLMGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGL 120  
Db 61 TMMHMSGSGGGLMGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGL 120  
Oy 121 TSTTNSPLDQGLNETSNDSTSTSTSTSSDPMQLKRESEINSLFDDQDQT 180  
Oy 121 TSTTNSPLDQGLNETSNDSTSTSTSTSSDPMQLKRESEINSLFDDQDQT 180  
Oy 181 QSSSSGCKPTFGDQWYKCVTDAISGLWNLGSLGGLGGLGGLGGLGGLGGL 240  
Db 181 QSSSSGCKPTFGDQWYKCVTDAISGLWNLGSLGGLGGLGGLGGLGGLGGL 240  
Oy 241 GKGGLNLSPYVQQLGNAVGTGIGKAGIQLNDGTHRRSTSTSPYKGDPMAMEI 300  
Oy 301 GQPMQDTPVYKQVQKQGVKVDKSNMAALSFPDDQDQTFASMEQPKAKOMIKR 360  
Oy 301 GQPMQDTPVYKQVQKQGVKVDKSNMAALSFPDDQDQTFASMEQPKAKOMIKR 360  
Oy 361 PMAGDTGNCNQAAGSGSLGIDAMAGDAINNALGKGA 403  
Db 361 PMAGDTGNCNQAAGSGSLGIDAMAGDAINNALGKGA 403

## RESULT 15

US-09-880-371-3  
! SOFTWARE: PatentIn Ver. 2.1  
! SEQ ID NO 3  
! TYPE: PRT  
! ORGANISM: Erwinia amylovora  
! US-09-880-371-3  
! APPLICATION US/9980371  
! INVENTOR: Wei, Zhong-Min  
! APPLICANT: Dabocher, Jay  
! TITLE OF INVENTION: PLANTS  
! FILE REFERENCE: 218/93/91  
! CURRENT FILING DATE: 2001-06-13  
! PRIOR APPLICATION NUMBER: 60/211,585  
! NUMBER OF SEQ ID NOS: 16  
! SOFTWARE: PatentIn Ver. 2.1  
! SEQ ID NO 3  
! TYPE: PRT  
! ORGANISM: Erwinia amylovora  
! US-09-880-371-3

Query Match 100.0% Score 2079; Db 22; Length 403;  
Best Local Similarity 100.0%; Pos 21; Indels 0; Gaps 0;  
Matches 403; Conservative 0; Mismatches 0;

Oy 1 NSLWTSGLASTPMQISIGAGNNGLLGTSRQNGALGAGGNGNTVWLAGLL 60  
Oy 1 NSLWTSGLASTPMQISIGAGNNGLLGTSRQNGALGAGGNGNTVWLAGLL 60  
Oy 61 TMMHMSGSGGGLMGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGL 120

Db 61 TMMHMSGSGGGLMGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGL 120  
Oy 121 TSTTNSPLDQGLNETSNDSTSTSTSSDPMQLKRESEINSLFDDQDQT 180  
Db 121 TSTTNSPLDQGLNETSNDSTSTSTSSDPMQLKRESEINSLFDDQDQT 180  
Oy 181 QSSSSGCKPTFGDQWYKCVTDAISGLWNLGSLGGLGGLGGLGGLGGLGGL 240  
Db 181 QSSSSGCKPTFGDQWYKCVTDAISGLWNLGSLGGLGGLGGLGGLGGLGGL 240  
Oy 241 GKGGLNLSPYVQQLGNAVGTGIGKAGIQLNDGTHRRSTSTSPYKGDPMAMEI 300  
Db 241 GKGGLNLSPYVQQLGNAVGTGIGKAGIQLNDGTHRRSTSTSPYKGDPMAMEI 300  
Oy 301 GQPMQDTPVYKQVQKQGVKVDKSNMAALSFPDDQDQTFASMEQPKAKOMIKR 360  
Db 301 GQPMQDTPVYKQVQKQGVKVDKSNMAALSFPDDQDQTFASMEQPKAKOMIKR 360  
Oy 361 PMAGDTGNCNQAAGSGSLGIDAMAGDAINNALGKGA 403  
Db 361 PMAGDTGNCNQAAGSGSLGIDAMAGDAINNALGKGA 403

Search completed: May 2, 2002, 03:35:07  
Job time: 4.18 sec



us-09-770-693-3.rapm

Thu May 2 11:45:17 2002







```

121 TRKAGNAYVCSQVGVHSICAMETSGGKIFCSGGGLG---GGGSGFGLATVWVNGP 177
Db 104 DMIGSS---LVATGSS---KCGN---TTFTHRLDNLGALG---INSTQNDGST---SCT 148
Db 178 UNKNGSLQNLGGLGGGAGNACTGGLSSGKLGKGLNLSQPVYQVQUGNAVGTGCM 267
Qy 149 DSTFSSDSQMKQLLWKFSEIQLPLFCDQDQGTGGSSGGKQPTFEDONAYKGYVTLASG 208
Db 238 GCGSSNS-----GGGSGSGSGSSGSS---SNDKN-----NCSGSG 270
Qy 209 LMKNLISQLNGKGLGGGAGNACTGGLSSGKLGKGLNLSQPVYQVQUGNAVGTGCM 267
Db 271 GSSGSGSGSSGSGSSGSGSGSSGSGS---SNDKREKSKENLLG----- 373
Qy 268 -----KAGLNADGTHRHSTFSTFYNGCDAMAKELQDPMQVPEVF 311
Db 330 HFKCQTPKGNRAGSGESGIGQPGQVY---SNNREKSKENLLG----- 373
Qy 312 GRYQYQNGQPVYTKDQSN 331
Db 374 -----GSDNTRQCGSSN 386

```

```

RESULT 5
US-10-006-117A-52
: SEQUENCE 52, Application US/10006117A
: GENERAL INFORMATION:
: APPLICANT: Baker, Kevin P.
: APPLICANT: Botstein, David
: APPLICANT: Benayahu, Yacov
: APPLICANT: Eron, Dan I.
: APPLICANT: Ferrata, Neapolone
: APPLICANT: Fong, Sherman
: APPLICANT: Goddard, Audrey
: APPLICANT: Godowski, Paul J.
: APPLICANT: Gurney, Austin L.
: APPLICANT: Hillan, Kenneth J.
: APPLICANT: Paoel, Nicholas F.
: TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
: ACID SEQUENCES Encoding the Same
: FILE REFERENCE: P2830747
: CURRENT APPLICATION NUMBER: US/00/06.117A
: CURRENT FILING DATE: 2001-03-19
: PRIOR FILING DATE: 2001-07-09
: NUMBER OF SEQ ID NOS: 477
: SEQ ID NO 1
: SEQ ID NO 2
: SEQ ID NO 3
: TYPE: PRT
: ORGANISM: Homo sapiens
US-10-006-117A-52
: SEQUENCE 52, Application US/10006117A
: GENERAL INFORMATION:
: APPLICANT: Baker, Kevin P.
: APPLICANT: Botstein, David
: APPLICANT: Benayahu, Yacov
: APPLICANT: Eron, Dan I.
: APPLICANT: Ferrata, Neapolone
: APPLICANT: Fong, Sherman
: APPLICANT: Goddard, Audrey
: APPLICANT: Godowski, Paul J.
: APPLICANT: Gurney, Austin L.
: APPLICANT: Hillan, Kenneth J.
: APPLICANT: Paoel, Nicholas F.
: TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
: ACID SEQUENCES Encoding the Same
: FILE REFERENCE: P2830747
: CURRENT APPLICATION NUMBER: US/00/06.117A
: CURRENT FILING DATE: 2001-03-19
: PRIOR FILING DATE: 2001-07-09
: NUMBER OF SEQ ID NOS: 477
: SEQ ID NO 1
: SEQ ID NO 2
: SEQ ID NO 3
: TYPE: PRT
: ORGANISM: Homo sapiens
US-10-006-117A-52

```

```

Query Match
Best Local Similarity 23.9%, Pred. No. 3.4e-06;
Matches 91; Conservative 37; Mismatches 141; Indels 111; Gaps 18;

Qy 9 GASTWVLSIGCGAGNGLIGTS---RNNGLGNSALG-----LG-----GQNDGT 52
Db 61 GAGSGVGLATGQVTRVATVGVFGAGLNAGHRYGCAHALNGLHETIGRADEV 120
Qy 53 VIOGLAFLTDKNH-----HNSMGCGGLG---GGLGAGLGGSGGLGELSHALN 103
Db 121 TRKAGNAYVCSQVGVHSICAMETSGGKIFCSGGGLG---GGGSGFGLATVWVNGP 177
Qy 104 DMIGSS---LVATGSS---KCGN---TTFTHRLDNLGALG---INSTQNDGST---SCT 148
Db 178 UNKNGSLQNLGGLGGGAGNACTGGLSSGKLGKGLNLSQPVYQVQUGNAVGTGCM 267
Qy 149 DSTFSSDSQMKQLLWKFSEIQLPLFCDQDQGTGGSSGGKQPTFEDONAYKGYVTLASG 208
Db 238 GCGSSNS-----GGGSGSGSGSSGSS---SNDKN-----NCSGSG 270
Qy 209 LMKNLISQLNGKGLGGGAGNACTGGLSSGKLGKGLNLSQPVYQVQUGNAVGTGCM 267
Db 271 GSSGSGSGSSGSGSGSGSGSSGSGS---SNDKREKSKENLLG----- 373

```

```

Qy 149 DSTFSSDSQMKQLLWKFSEIQLPLFCDQDQGTGGSSGGKQPTFEDONAYKGYVTLASG 208
Db 238 GCGSSNS-----GGGSGSGSGSGSS---SNDKN-----NCSGSG 270
Qy 209 LMKNLISQLNGKGLGGGAGNACTGGLSSGKLGKGLNLSQPVYQVQUGNAVGTGCM 267
Db 271 GSSGSGSGSSGSGSGSGSGSGSSGSGS---SNDKREKSKENLLG----- 373
Qy 268 -----KAGLNADGTHRHSTFSTFYNGCDAMAKELQDPMQVPEVF 311
Db 330 HFKCQTPKGNRAGSGESGIGQPGQVY---SNNREKSKENLLG----- 373
Qy 312 GRYQYQNGQPVYTKDQSN 331
Db 374 -----GSDNTRQCGSSN 386

```

```

RESULT 6
US-10-006-130A-52
: SEQUENCE 6, Application US/10006130A
: GENERAL INFORMATION:
: APPLICANT: Baker, Kevin P.
: APPLICANT: Botstein, David
: APPLICANT: Benayahu, Yacov
: APPLICANT: Eron, Dan I.
: APPLICANT: Ferrata, Neapolone
: APPLICANT: Fong, Sherman
: APPLICANT: Goddard, Audrey
: APPLICANT: Godowski, Paul J.
: APPLICANT: Gurney, Austin L.
: APPLICANT: Hillan, Kenneth J.
: APPLICANT: Paoel, Nicholas F.
: TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
: ACID SEQUENCES Encoding the Same
: FILE REFERENCE: P2830747
: CURRENT APPLICATION NUMBER: US/00/06.130A
: CURRENT FILING DATE: 2001-03-19
: PRIOR APPLICATION REMOVED - See File Wrapper or Palm
: NUMBER OF SEQ ID NOS: 477
: SEQ ID NO 52
: SEQ ID NO 53
: SEQ ID NO 54
: TYPE: PRT
: ORGANISM: Homo sapiens
US-10-006-130A-52

```

```

Query Match
Best Local Similarity 23.9%, Pred. No. 3.4e-06;
Matches 91; Conservative 37; Mismatches 141; Indels 111; Gaps 18;

Qy 9 GASTWVLSIGCGAGNGLIGTS---RNNGLGNSALG-----LG-----GQNDGT 52
Db 61 GAGSGVGLATGQVTRVATVGVFGAGLNAGHRYGCAHALNGLHETIGRADEV 120
Qy 53 VIOGLAFLTDKNH-----HNSMGCGGLG---GGLGAGLGGSGGLGELSHALN 103
Db 121 TRKAGNAYVCSQVGVHSICAMETSGGKIFCSGGGLG---GGGSGFGLATVWVNGP 177
Qy 104 DMIGSS---LVATGSS---KCGN---TTFTHRLDNLGALG---INSTQNDGST---SCT 148
Db 178 UNKNGSLQNLGGLGGGAGNACTGGLSSGKLGKGLNLSQPVYQVQUGNAVGTGCM 267
Qy 149 DSTFSSDSQMKQLLWKFSEIQLPLFCDQDQGTGGSSGGKQPTFEDONAYKGYVTLASG 208
Db 238 GCGSSNS-----GGGSGSGSGSSGSS---SNDKN-----NCSGSG 270
Qy 209 LMKNLISQLNGKGLGGGAGNACTGGLSSGKLGKGLNLSQPVYQVQUGNAVGTGCM 267
Db 271 GSSGSGSGSSGSGSGSGSGSGSGS---SNDKREKSKENLLG----- 373

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07 268 -----FACTQALMDJOTHRHSTFRVYKNDKAMKEIOPNDQYPRV 311
DB 330 HNGCKCFEFGNARSGSSHQFROGVY--SSNMHEISEKRNELLO----- 373
07 312 GKPYOKGPGQDVKDYSK 331
DB 374 -----GSDJNRQGGSSW 386

RESULT
Sequence 7 129-53
Sequence 52 Application US/1006172A
GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Baker, Kevin P.
APPLICANT: Debnoyers, Luc
APPLICANT: Eaton, Dan L.
APPLICANT: Fournier, Jean-Pierre
APPLICANT: Fournier, Jean-Pierre
APPLICANT: Gao, Wei-Qiang
APPLICANT: Goddard, Audrey
APPLICANT: Goddard, Audrey
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Gurney, Austin L.
APPLICANT: Hildesheim, Kenneth J.
APPLICANT: Pan, James
APPLICANT: Pan, Nicholas F.
INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: P28302111 According the same
FILE REFERENCE: P28302111 According the same
CURRENT APPLICATION NUMBER: US/10/006,172A
PRIORITY APPLICATION NUMBER: 60/098716
PRIOR FILING DATE: 1998-09-01
PRIOR FILING DATE: 1998-09-01
PRIOR FILING DATE: 1998-09-01
PRIOR APPLICATION NUMBER: 60/098749
PRIOR FILING DATE: 1998-09-01
PRIOR FILING DATE: 1998-09-01
PRIOR APPLICATION NUMBER: 60/098750
PRIOR FILING DATE: 1998-09-01
PRIOR APPLICATION NUMBER: 60/098803
PRIOR FILING DATE: 1998-09-01
PRIOR APPLICATION NUMBER: 60/098821
PRIOR FILING DATE: 1998-09-02
PRIOR FILING DATE: 1998-09-02
PRIOR APPLICATION NUMBER: 60/098843
PRIOR FILING DATE: 1998-09-02
PRIOR APPLICATION NUMBER: 60/098936
PRIOR FILING DATE: 1998-09-03
PRIOR APPLICATION NUMBER: 60/098956
PRIOR FILING DATE: 1998-09-03
PRIOR APPLICATION NUMBER: 60/098958
PRIOR FILING DATE: 1998-09-03
PRIOR APPLICATION NUMBER: 60/099602
PRIOR FILING DATE: 1998-09-09
PRIOR APPLICATION NUMBER: 60/099642
PRIOR FILING DATE: 1998-09-10
PRIOR APPLICATION NUMBER: 60/099741
PRIOR FILING DATE: 1998-09-10
PRIOR APPLICATION NUMBER: 60/099754
PRIOR FILING DATE: 1998-09-10
PRIOR APPLICATION NUMBER: 60/099763
PRIOR FILING DATE: 1998-09-10
PRIOR APPLICATION NUMBER: 60/099792
PRIOR FILING DATE: 1998-09-10
PRIOR APPLICATION NUMBER: 60/099808
PRIOR FILING DATE: 1998-09-10
PRIOR APPLICATION NUMBER: 60/099812
PRIOR FILING DATE: 1998-09-10
PRIOR APPLICATION NUMBER: 60/099815
PRIOR FILING DATE: 1998-09-10
PRIOR APPLICATION NUMBER: 60/099816
PRIOR FILING DATE: 1998-09-10
PRIOR APPLICATION NUMBER: 60/040385
PRIOR FILING DATE: 1998-09-15
PRIOR APPLICATION NUMBER: 60/100388
PRIOR FILING DATE: 1998-09-15
PRIOR APPLICATION NUMBER: 60/100390
PRIOR FILING DATE: 1998-09-15
PRIOR APPLICATION NUMBER: 60/100584
PRIOR FILING DATE: 1998-09-15
PRIOR APPLICATION NUMBER: 60/100627
PRIOR FILING DATE: 1998-09-15
PRIOR APPLICATION NUMBER: 60/100661
PRIOR FILING DATE: 1998-09-15
PRIOR APPLICATION NUMBER: 60/100662
PRIOR FILING DATE: 1998-09-15
PRIOR APPLICATION NUMBER: 60/100684
PRIOR FILING DATE: 1998-09-15
PRIOR APPLICATION NUMBER: 60/100683
PRIOR FILING DATE: 1998-09-17
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PRIOR APPLICATION NUMBER: 60/100710
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PRIOR FILING DATE: 1998-09-17
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PRIOR FILING DATE: 1998-09-18
PRIOR APPLICATION NUMBER: 60/100849
PRIOR FILING DATE: 1998-09-17
PRIOR APPLICATION NUMBER: 60/100919
PRIOR FILING DATE: 1998-09-17
PRIOR APPLICATION NUMBER: 60/100930
PRIOR FILING DATE: 1998-09-17
PRIOR APPLICATION NUMBER: 60/101014
PRIOR FILING DATE: 1998-09-18
PRIOR APPLICATION NUMBER: 60/101068
PRIOR FILING DATE: 1998-09-18
PRIOR APPLICATION NUMBER: 60/101071
PRIOR FILING DATE: 1998-09-22
PRIOR APPLICATION NUMBER: 60/101279
PRIOR FILING DATE: 1998-09-22
PRIOR APPLICATION NUMBER: 60/101471
PRIOR FILING DATE: 1998-09-23
PRIOR APPLICATION NUMBER: 60/101472
PRIOR FILING DATE: 1998-09-23
PRIOR APPLICATION NUMBER: 60/101474
PRIOR FILING DATE: 1998-09-23
PRIOR APPLICATION NUMBER: 60/101475
PRIOR FILING DATE: 1998-09-23
PRIOR APPLICATION NUMBER: 60/101476
PRIOR FILING DATE: 1998-09-23
PRIOR APPLICATION NUMBER: 60/101477
PRIOR FILING DATE: 1998-09-23
PRIOR APPLICATION NUMBER: 60/101479
PRIOR FILING DATE: 1998-09-24
PRIOR APPLICATION NUMBER: 60/101741
PRIOR FILING DATE: 1998-09-24
PRIOR APPLICATION NUMBER: 60/101743
PRIOR FILING DATE: 1998-09-24
PRIOR APPLICATION NUMBER: 60/101915
PRIOR FILING DATE: 1998-09-24
PRIOR APPLICATION NUMBER: 60/101916
PRIOR FILING DATE: 1998-09-25
PRIOR APPLICATION NUMBER: 60/102207
PRIOR FILING DATE: 1998-09-25
PRIOR APPLICATION NUMBER: 60/102240
PRIOR FILING DATE: 1998-09-25
PRIOR APPLICATION NUMBER: 60/102307
PRIOR FILING DATE: 1998-09-25
PRIOR APPLICATION NUMBER: 60/102330
PRIOR FILING DATE: 1998-09-29
PRIOR APPLICATION NUMBER: 60/102331
PRIOR FILING DATE: 1998-09-29

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[illegible]

PRIOR FILING DATE: 1998-10-21  
 PRIOR FILING DATE: 1998-10-21  
 PRIOR FILING DATE: 1998-10-22  
 PRIOR FILING DATE: 1998-10-22  
 PRIOR FILING DATE: 1998-10-26  
 PRIOR FILING DATE: 1998-10-26  
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 PRIOR FILING DATE: 1998-10-27  
 PRIOR FILING DATE: 1998-10-27  
 PRIOR FILING DATE: 1998-10-27  
 PRIOR FILING DATE: 1998-10-28  
 PRIOR FILING DATE: 1998-10-28

PRIOR APPLICATION NUMBER: 60/105169  
 PRIOR APPLICATION NUMBER: 60/105169  
 PRIOR APPLICATION NUMBER: 60/105266  
 PRIOR APPLICATION NUMBER: 60/105693  
 PRIOR APPLICATION NUMBER: 60/105694  
 PRIOR APPLICATION NUMBER: 60/105694  
 PRIOR APPLICATION NUMBER: 60/105807  
 PRIOR APPLICATION NUMBER: 60/105882  
 PRIOR APPLICATION NUMBER: 60/105881  
 PRIOR APPLICATION NUMBER: 60/106023  
 PRIOR APPLICATION NUMBER: 60/106023  
 PRIOR APPLICATION NUMBER: 60/106029

Query Match  
 Best Local Similarity 23.9%; Pred. No. 46-06; Length 440;  
 Matches 91; Conservative 37; Mismatches 141; Indels 111; Gaps 18;

QY 9 GASTFQISQGGGNGGLGTS-RQNGAGLNSALG-----LG-----GQGMQNT 52  
 DB 61 GAGGYSVEALGGTREAIVGVYGVFGAALGNRYGVBAHALGTHGIEGRQAEV 120  
 QY 53 VQNLGALGLTGMN-----HMSMGSGGLMS- GELSGGLGNGELSGGELGRLSIAL 103  
 DB 121 TRHGAIVRSGWVPHSGAMETSGRIGSGGGLG-----GQGMQNTGLTPWHPV 177  
 QY 104 DKLGGS-LVITGS-KGWN-----TFTSTNPIQALG-INSTNDQST---SQT 148  
 DB 178 GNSAGSGFPMQAPMGQMGNGPMPGINTGVAPQVGYSVRYNBNQDCTNPPFSG 237  
 QY 149 DSTSSHSPWQLLKWFSMTSGFQDQDQVSGSSGKQPTGDMATKGVITPV 208  
 DB 238 GQSSGSS-----GGSGSGSGSGSGS--SNGDN-----NWSGG 270  
 QY 201 LKNGLSLQLGLGGGGGAGTGLDGLSLGKGLNGLGVYQLANVGTGN- 267  
 DB 271 GSSGSSGSSGSSGSSGSSGSGS--SNGSGSGSGSGSGSWSSTVSNGSGGGG 329  
 QY 268 -----FACIQALNDITGRHSSTFVYKNGDAMAKETGQMPQVFPV 311  
 DB 330 HKPCKEPKNGRSGSSTGFGPGQV--SNNRISKKNLLG----- 373  
 QY 312 GPKQYKGPQVYTDKSN 331  
 DB 374 -----GSDINTRGSGSN 386

RESULT 12  
 US-10-006-041A-52  
 Sequence 52, Application US/1006618A  
 APPLICANT: Baker, Kevin P.  
 APPLICANT: Botstein, David  
 APPLICANT: Denoyers, Luc  
 APPLICANT: Balogh, Dan L.  
 APPLICANT: Ferrara, Napoleone  
 APPLICANT: Fong, Sherman  
 APPLICANT: Fong, Sherman  
 APPLICANT: Goddard, Audrey  
 APPLICANT: Godowski, Paul J.  
 APPLICANT: Grimaldi, Christopher J.  
 APPLICANT: Hillan, Kenneth J.  
 APPLICANT: Hillan, Kenneth J.  
 APPLICANT: Pan, James  
 APPLICANT: Pan, James  
 TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic  
 TITLE OF INVENTION: Acids Encoding the Same  
 CURRENT APPLICATION NUMBER: US/10/006.818A  
 PRIOR FILING DATE: 2001-12-06  
 PRIOR APPLICATION REMOVED - See File Wrapper or Palm  
 SEQ ID NO 52  
 LENGTH: 440  
 ORGANISM: Homo sapiens

PRIOR FILING DATE: 1998-10-21  
 PRIOR FILING DATE: 1998-10-21  
 PRIOR FILING DATE: 1998-10-22  
 PRIOR FILING DATE: 1998-10-22  
 PRIOR FILING DATE: 1998-10-26  
 PRIOR FILING DATE: 1998-10-26  
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 PRIOR FILING DATE: 1998-10-27  
 PRIOR FILING DATE: 1998-10-27  
 PRIOR FILING DATE: 1998-10-27  
 PRIOR FILING DATE: 1998-10-28  
 PRIOR FILING DATE: 1998-10-28

PRIOR APPLICATION NUMBER: 60/105169  
 PRIOR APPLICATION NUMBER: 60/105169  
 PRIOR APPLICATION NUMBER: 60/105266  
 PRIOR APPLICATION NUMBER: 60/105693  
 PRIOR APPLICATION NUMBER: 60/105694  
 PRIOR APPLICATION NUMBER: 60/105694  
 PRIOR APPLICATION NUMBER: 60/105807  
 PRIOR APPLICATION NUMBER: 60/105882  
 PRIOR APPLICATION NUMBER: 60/105881  
 PRIOR APPLICATION NUMBER: 60/106023  
 PRIOR APPLICATION NUMBER: 60/106023  
 PRIOR APPLICATION NUMBER: 60/106029

Query Match  
 Best Local Similarity 23.9%; Pred. No. 37-06; Length 440;  
 Matches 91; Conservative 37; Mismatches 141; Indels 111; Gaps 18;

QY 9 GASTFQISQGGGNGGLGTS-RQNGAGLNSALG-----LG-----GQGMQNT 52  
 DB 61 GAGGYSVEALGGTREAIVGVYGVFGAALGNRYGVBAHALGTHGIEGRQAEV 120  
 QY 53 VQNLGALGLTGMN-----HMSMGSGGLMS- GELSGGLGNGELSGGELGRLSIAL 103  
 DB 121 TRHGAIVRSGWVPHSGAMETSGRIGSGGGLG-----GQGMQNTGLTPWHPV 177  
 QY 104 DKLGGS-LVITGS-KGWN-----TFTSTNPIQALG-INSTNDQST---SQT 148  
 DB 178 GNSAGSGFPMQAPMGQMGNGPMPGINTGVAPQVGYSVRYNBNQDCTNPPFSG 237  
 QY 149 DSTSSHSPWQLLKWFSMTSGFQDQDQVSGSSGKQPTGDMATKGVITPV 208  
 DB 238 GQSSGSS-----GGSGSGSGSGSGS--SNGDN-----NWSGG 270  
 QY 201 LKNGLSLQLGLGGGGGAGTGLDGLSLGKGLNGLGVYQLANVGTGN- 267  
 DB 271 GSSGSSGSSGSSGSSGSSGSGS--SNGSGSGSGSGSGSWSSTVSNGSGGGG 329  
 QY 268 -----FACIQALNDITGRHSSTFVYKNGDAMAKETGQMPQVFPV 311  
 DB 330 HKPCKEPKNGRSGSSTGFGPGQV--SNNRISKKNLLG----- 373  
 QY 312 GPKQYKGPQVYTDKSN 331  
 DB 374 -----GSDINTRGSGSN 386

RESULT 11  
 US-10-006-041A-52  
 Sequence 52, Application US/1006618A  
 APPLICANT: Baker, Kevin P.  
 APPLICANT: Botstein, David  
 APPLICANT: Denoyers, Luc  
 APPLICANT: Balogh, Dan L.  
 APPLICANT: Ferrara, Napoleone  
 APPLICANT: Fong, Sherman  
 APPLICANT: Fong, Sherman  
 APPLICANT: Goddard, Audrey  
 APPLICANT: Godowski, Paul J.  
 APPLICANT: Grimaldi, Christopher J.  
 APPLICANT: Hillan, Kenneth J.  
 APPLICANT: Hillan, Kenneth J.  
 APPLICANT: Pan, James  
 APPLICANT: Pan, James  
 TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic  
 TITLE OF INVENTION: Acids Encoding the Same

US-10-006-818A-52

Query Match 8.48; Score 174.5; Db 6; Length 440;  
 Best Local Similarity 23.9%; Pred. No. 3.4e-06;  
 Matches 91; Conservative 37; Mismatches 141; Indels 111; Gaps 18;

Qy 9 GASTMISIGAGAGNGLLST-KRMGLGNSALG-----LG-----GKQKNT 52  
 Db 61 GAGCKSVGADQTBVAVGVTVPTGDAADLNNVRAHMAALGHTHEIGRQEDV 120  
 Qy 53 VNIQALLCTQNM-----MSHMGSGGLG-----GGLGGLGAGLGLGSLGSLN 103  
 Db 121 IHRGADVAIVGQVGVHSQWMTSGHGTGGGGGLG-----GQGNQNT 177  
 Qy 104 DMUGGS-LMTLGS-KGGIN-----TSTTNSPDALG-INFTSONDST---SOT 148  
 Db 178 GNSGSGFQMGQAGWQGNMGNGPMTGTVAGVQGVTSVAMNHCCTNFPSS 237  
 Qy 61 GAGCKSVGADQTBVAVGVTVPTGDAADLNNVRAHMAALGHTHEIGRQEDV 120  
 Qy 53 VNIQALLCTQNM-----MSHMGSGGLG-----GGLGGLGAGLGLGSLN 103  
 Db 121 IHRGADVAIVGQVGVHSQWMTSGHGTGGGGGLG-----GQGNQNT 177  
 Qy 104 DMUGGS-LMTLGS-KGGIN-----TSTTNSPDALG-INFTSONDST---SOT 148  
 Db 178 GNSGSGFQMGQAGWQGNMGNGPMTGTVAGVQGVTSVAMNHCCTNFPSS 237  
 Qy 149 DTSUSSDPMOQLKMFSLMGLFQDQVQVSSSGQVGFTRQWNTKRYATLSS 208  
 Db 238 GGGSSNS-----GGSSGSGSSGSS-----SNDINN-----NSSSG 270  
 Qy 209 LAMGLGQLAGLGLGAGNACTGAGLGLGAGLGLGAGLGLGAGLGLGAGLGL 267  
 Db 271 GSS 329  
 Qy 268 -----FACIADNDGTHIRSTSTFYVKGDKRAKETIQMOTPEV 311  
 Db 330 HPCQKCFQKFAKSGSSTQGFQGV---SNNRSLSKNRLGLG----- 373  
 Qy 312 GFKYQKQGVGVWYDKSN 331  
 Db 374 -----GSDNYFGSGSM 386

RESULT 13

US-10-015-386A-52  
 : Sequence 52, Application US/1001212A

GENERAL INFORMATION:  
 : TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic  
 : ACIDS Encoding the Same  
 : APPLICANT: Baker, Kevin P.  
 : APPLICANT: Betstein, David  
 : APPLICANT: Desoyers, Luc  
 : APPLICANT: Faton, Ian J.  
 : APPLICANT: Fong, Sherman  
 : APPLICANT: Gao, Wei-Qiang  
 : APPLICANT: Godowski, Paul J.  
 : APPLICANT: Grimaldi, Christopher J.  
 : APPLICANT: Hillan, Kenneth J.  
 : APPLICANT: Hillan, Kenneth J.  
 : APPLICANT: Paoi, Nicholas F.  
 : TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic  
 : ACIDS Encoding the Same  
 : CURRENT APPLICATION NUMBER: US/012/121A  
 : CURRENT FILING DATE: 2001-12-07  
 : NUMBER OF SEQ ID NOS: 477  
 : SEQ ID NO 52 ID NOS: 477  
 : SEQ ID NO 52 ID NOS: 477  
 : TYPE: PAT  
 : ORGANISM: Homo sapiens  
 : US-10-012-121A-52

Query Match 8.48; Score 174.5; Db 6; Length 440;  
 Best Local Similarity 23.9%; Pred. No. 3.4e-06;  
 Matches 91; Conservative 37; Mismatches 141; Indels 111; Gaps 18;

Qy 9 GASTMISIGAGAGNGLLST-KRMGLGNSALG-----LG-----GKQKNT 52

Db 61 GAGCKSVGADQTBVAVGVTVPTGDAADLNNVRAHMAALGHTHEIGRQEDV 120  
 Qy 53 VNIQALLCTQNM-----MSHMGSGGLG-----GGLGGLGAGLGLGSLN 103  
 Db 121 IHRGADVAIVGQVGVHSQWMTSGHGTGGGGGLG-----GQGNQNT 177  
 Qy 104 DMUGGS-LMTLGS-KGGIN-----TSTTNSPDALG-INFTSONDST---SOT 148  
 Db 178 GNSGSGFQMGQAGWQGNMGNGPMTGTVAGVQGVTSVAMNHCCTNFPSS 237  
 Qy 149 DTSUSSDPMOQLKMFSLMGLFQDQVQVSSSGQVGFTRQWNTKRYATLSS 208  
 Db 238 GGGSSNS-----GGSSGSGSSGSS-----SNDINN-----NSSSG 270  
 Qy 209 LAMGLGQLAGLGLGAGNACTGAGLGLGAGLGLGAGLGLGAGLGLGAGLGL 267  
 Db 271 GSS 329  
 Qy 268 -----FACIADNDGTHIRSTSTFYVKGDKRAKETIQMOTPEV 311  
 Db 330 HPCQKCFQKFAKSGSSTQGFQGV---SNNRSLSKNRLGLG----- 373  
 Qy 312 GFKYQKQGVGVWYDKSN 331  
 Db 374 -----GSDNYFGSGSM 386

RESULT 14

US-10-015-386A-52  
 : Sequence 52, Application US/10015386A

GENERAL INFORMATION:  
 : TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic  
 : ACIDS Encoding the Same  
 : APPLICANT: Baker, Kevin P.  
 : APPLICANT: Betstein, David  
 : APPLICANT: Desoyers, Luc  
 : APPLICANT: Faton, Ian J.  
 : APPLICANT: Fong, Sherman  
 : APPLICANT: Godowski, Paul J.  
 : APPLICANT: Grimaldi, Christopher J.  
 : APPLICANT: Hillan, Kenneth J.  
 : APPLICANT: Hillan, Kenneth J.  
 : APPLICANT: Paoi, Nicholas F.  
 : TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic  
 : ACIDS Encoding the Same  
 : CURRENT APPLICATION NUMBER: US/10/15.386A  
 : CURRENT FILING DATE: 2001-12-07  
 : NUMBER OF SEQ ID NOS: 477  
 : SEQ ID NO 52 ID NOS: 477  
 : SEQ ID NO 52 ID NOS: 477  
 : TYPE: PAT  
 : ORGANISM: Homo sapiens  
 : US-10-015-386A-52

Query Match 8.48; Score 174.5; Db 6; Length 440;  
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 Qy 53 VNIQALLCTQNM-----MSHMGSGGLG-----GGLGGLGAGLGLGSLN 103  
 Db 121 IHRGADVAIVGQVGVHSQWMTSGHGTGGGGGLG-----GQGNQNT 177  
 Qy 104 DMUGGS-LMTLGS-KGGIN-----TSTTNSPDALG-INFTSONDST---SOT 148

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Db 178 ONSGASFOHQAHPQGGGGGPPHRTGTTGAVQAQTGVSVASNSNGCSTNPPFSS 337
Oy 149 DSTDSGSMQGLLAWSEWELKSLFQSDQDTGSSGCKPTGDEBNATKGYDALSG 208
Db 238 GGSNSN-----GGSGSGSGSGS---SNDNR-----NSSSG 270
Oy 209 LKNGELSGLLGCGCGGAGNAGTGLSSGCKGLGNTSPDYVLGHVAVGTGK- 267
Db 271 GSSGSGSGSGSGSGSGSGSGS-SGNSGSGSGSGSGSGSGSGSGSGSGSGSG 329
Oy 268 -----KAGTALNDLQTHRHSTSFYKNGGDMAKELGQPMOYPIEVF 311
Db 330 HRFQCEKPFHARGSGSGTGFQGGV---SSNMREISKENLLG----- 373
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Db 374 -----GSDNFRGQGSN 386

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Search completed: May 2, 2002, 03:36:03  
Job Time: 268 sec

## RESULTS

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US-10-015-387A-52
> Sequence 52, Application US/A0015387A
> GENERAL INFORMATION:
> APPLICANT: Botstein, David P.
> APPLICANT: Botstein, David
> APPLICANT: Denoyers, Luc
> APPLICANT: Denoyers, Luc
> APPLICANT: Farrara, Napoleone
> APPLICANT: Fong, Sherman
> APPLICANT: Goddard, Audrey
> APPLICANT: Goddard, Paul J.
> APPLICANT: Grimaldi, Christopher J.
> APPLICANT: Hillan, Kenneth J.
> APPLICANT: Pan, James
> APPLICANT: Pan, James
> APPLICANT: Pan, James
> TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
> ACIDS ENCODING THE SAME
> CURRENT APPLICATION NUMBER: US/10/015 387A
> PRIOR APPLICATION NUMBER: 2001-12-12
> Prior Application removed - See File Wrapper or Palm
> SEQ ID NO 52
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> ORGANISM: Homo sapiens
> ORGANISM: Homo sapiens
US-10-015-387A-52

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Query Match 8.44; Score 174.5; Db 6; Length 440;
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Oy 104 DKGGS--LNTLS--KGNH-----TETINSELDALG--INFTNDSST---SGT 148
Db 178 ONSGASFOHQAHPQGGGGGPPHRTGTTGAVQAQTGVSVASNSNGCSTNPPFSS 237
Oy 149 DSTDSGSMQGLLAWSEWELKSLFQSDQDTGSSGCKPTGDEBNATKGYDALSG 208
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GenCore version 4.5  
Copyright (c) 1993 - 2000 CompuGen Ltd.

OM nucleic - nucleic search, using sw model

Run on: May 2, 2002, 01:37:15 ; Search time 4673.21 Seconds  
(without alignments)  
7596.017 Million cell updates/sec

Title: US-09-770-693-4

Perfect score: 1288

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Scoring table: IDENTITY\_NUC

Gapop 10.0 , Gapext 1.0

Searched: 2203303 seqs, 1083143700 residues

Total number of hits satisfying chosen parameters: 4404606

Minimum DB seq length: 0

Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

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Frnd. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

# SUMMARIES

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? STRANDEDNESS: single
? TOPOLOGY: linear
? MOLECULE TYPE: cDNA
US-08-851-376-3

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? LENGTH: 41206
? TYPE: DNA
? STR: Homo sapiens
US-10-105-295-6862

Query Match          2.9%  Score 36.4; DB 6; Length 41206;
Best Local Similarity 51.9%; Pred. No. 2;
Matches 83; Conservative 0; Mismatches 77; Indels 0; Gaps 0;
Oy 688 ctgtcgccgtgagtgatgaatgacgcagctctctggaaaggggagctggaggt 747
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Db 40661 Cttggagccgcttggacacgacgagccctctggccgacacgacgagctgca 40662

Oy 748 agccggcggtgaagcgagcggtgctgcgcgagtcgcgtcgctggcgcaaaagctg 807
||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
Db 40601 GCAGCCAGCGGACGCCAGGAGGAGGAGGCTTGGAATGACGACGCTGCCAGGGGTAGG 40642

Oy 808 caaacacgcagggcgagggcgagctgacacgcagcttgcagaa 847
||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
Db 40541 CAAACACTGAGTGTCTCTGTGAGCCCTCGAATTGTCCA 05092

RESULT 3 2105-1976
US-09-540-2105-1976
? TITLE: DNA
? STR: Homo sapiens
? CURRENT FILING DATE: 2002-04-03
? GENERAL INFORMATION:
? APPLICANT: Seilheimer, Jeffrey J.
? APPLICANT: Stuart, Susan G.
? APPLICANT: Sluve, Laura L.
? APPLICANT: Nauplion, Rebecca E.
? TITLE OF INVENTION: POLYNUCLEOTIDES OF URINARY TRACT TISSUE
? CURRENT FILING DATE: 2002-04-03
? PRIOR FILING DATE: March 12, 1998
? PRIOR APPLICATION NUMBER: 08/78,199
? PRIOR FILING DATE: April 10, 1996
? PRIOR APPLICATION NUMBER: 08/395,244
? PRIOR FILING DATE: February 16, 1995
? PRIOR FILING DATE: September 27, 1996
? PRIOR APPLICATION NUMBER: 60/005,526
? PRIOR APPLICATION NUMBER: 08/824,025
? PRIOR FILING DATE: March 25, 1997
? PRIOR FILING DATE: March 25, 1996
? PRIOR FILING DATE: March 29, 1996
? PRIOR APPLICATION NUMBER: 08/826,847
? PRIOR APPLICATION NUMBER: 60/015,533
? PRIOR FILING DATE: April 10, 1996
? PRIOR FILING DATE: November 31, 1995
? PRIOR APPLICATION NUMBER: 60/023,308
? PRIOR FILING DATE: July 31, 1996
? PRIOR APPLICATION NUMBER: 08/78,178
? PRIOR FILING DATE: May 22, 1997
? PRIOR APPLICATION NUMBER: 60/018,217
? PRIOR FILING DATE: June 24, 1997
? PRIOR APPLICATION NUMBER: 08/881,589
? PRIOR FILING DATE: June 25, 1996
? PRIOR APPLICATION NUMBER: 08/802,402
? PRIOR FILING DATE: July 31, 1996
? PRIOR APPLICATION NUMBER: 60/023,308
? PRIOR FILING DATE: July 31, 1996
? PRIOR APPLICATION NUMBER: 08/78,981
? PRIOR FILING DATE: August 3, 1996
? PRIOR APPLICATION NUMBER: 60/025,041

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? PRIOR APPLICATION NUMBER: 08/903,471
? PRIOR FILING DATE: July 30, 1997
? PRIOR APPLICATION NUMBER: 60/026,578
? PRIOR FILING DATE: July 31, 1996
? PRIOR APPLICATION NUMBER: 08/903,556
? PRIOR FILING DATE: August 21, 1996
? PRIOR APPLICATION NUMBER: 60/025,217
? PRIOR FILING DATE: August 22, 1996
? PRIOR APPLICATION NUMBER: 08/937,183
? PRIOR APPLICATION NUMBER: 60/026,598
? PRIOR FILING DATE: September 24, 1996
? PRIOR FILING DATE: October 25, 1997
? PRIOR APPLICATION NUMBER: 60/030,144
? PRIOR APPLICATION NUMBER: 08/826,847
? PRIOR FILING DATE: April 10, 1997
? PRIOR APPLICATION NUMBER: 08/826,847
? PRIOR FILING DATE: April 10, 1996
? PRIOR APPLICATION NUMBER: 08/755,524
? PRIOR FILING DATE: November 22, 1995
? PRIOR APPLICATION NUMBER: 60/027,496
? PRIOR FILING DATE: November 22, 1995
? PRIOR APPLICATION NUMBER: 09/021,031
? PRIOR APPLICATION NUMBER: 60/039,325
? PRIOR FILING DATE: February 13, 1997
? PRIOR FILING DATE: March 4, 1998
? PRIOR APPLICATION NUMBER: 60/040,431
? PRIOR FILING DATE: March 12, 1998
? PRIOR APPLICATION NUMBER: 09/041,894
? PRIOR FILING DATE: March 14, 1997
? PRIOR APPLICATION NUMBER: 09/050,817
? PRIOR FILING DATE: March 30, 1997
? PRIOR APPLICATION NUMBER: 60/047,792
? PRIOR FILING DATE: April 11, 1997
? PRIOR APPLICATION NUMBER: 09/094,999
? PRIOR APPLICATION NUMBER: 60/048,431
? PRIOR FILING DATE: May 29, 1997
? PRIOR FILING DATE: June 30, 1998
? PRIOR APPLICATION NUMBER: 60/052,751
? PRIOR APPLICATION NUMBER: 09/034,079
? PRIOR FILING DATE: June 9, 1998
? PRIOR APPLICATION NUMBER: 60/048,975
? PRIOR FILING DATE: June 13, 1997
? NUMBER OF SEQ ID NOS: 35654
? OPTIMUM BASE PAIR PROGRAM
? SEQ LENGTH: 15876
? TYPE: DNA
? STR: Homo sapiens
? NAME/REV: misc.feature
? FEATURE:
US-09-540-2108-15876
Query Match          2.8%  Score 36.4; DB 5; Length 262;
Best Local Similarity 56.8%; Pred. No. 0.42;
Matches 67; Conservative 0; Mismatches 51; Indels 0; Gaps 0;
Oy 228 cctcacgcgcgtgctgctaccacgcagatgatgatgatgatgatgatgatgag 287
||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
Db 40 ctctacacgttgctacacacacacacacacacacacacacacacacacacacac 99

Oy 288 tgcgtgagcgtgatgcggtgcgtgcgtgcgtgcgtgcgtgcgtgcgtgcgtgc 345

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Genbank Version 4.5  
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OM protein - protein search, using sw model

Run on: May 2, 2002, 03:21:24 : Search time 28.26 seconds

1055.571 Million cell updates/aec

Title: us-09-770-693-3

Perfect score: 2079

Sequence: I HSLVYSGIDALQVHSLQCA.....DAWMDKIMNNLQKLM 403

Scoring table:

Gap: 10.0, Gap: 0.5

Search: 522463 seqs, 74071320 residues

Total number of hits satisfying chosen parameters: 522463

Minimum db seq length: 0

Maximum db seq length: 200000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Display first 45 summaries

Databases:

1: /SI02/gqdata/geneseq/geneseq/AA1980.DMT.\*  
2: /SI02/gqdata/geneseq/geneseq/AA1981.DMT.\*  
3: /SI02/gqdata/geneseq/geneseq/AA1982.DMT.\*  
4: /SI02/gqdata/geneseq/geneseq/AA1983.DMT.\*  
5: /SI02/gqdata/geneseq/geneseq/AA1984.DMT.\*  
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7: /SI02/gqdata/geneseq/geneseq/AA1986.DMT.\*  
8: /SI02/gqdata/geneseq/geneseq/AA1987.DMT.\*  
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18: /SI02/gqdata/geneseq/geneseq/AA1997.DMT.\*  
19: /SI02/gqdata/geneseq/geneseq/AA1998.DMT.\*  
20: /SI02/gqdata/geneseq/geneseq/AA1999.DMT.\*  
21: /SI02/gqdata/geneseq/geneseq/AA2000.DMT.\*  
22: /SI02/gqdata/geneseq/geneseq/AA2001.DMT.\*

prod. No. is the number of results produced by change to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Length	DB ID	Description
1	2079	100.0	403	EWLnia amylovera
2	2079	100.0	403	Hyperresistive res
3	2079	100.0	403	Hyperresistive res
4	2079	100.0	403	Hyperresistive res
5	2079	100.0	403	Hyperresistive res
6	2079	100.0	403	Hyperresistive res
7	2079	100.0	403	Hyperresistive res
8	2079	100.0	403	Hyperresistive res
9	1933	92.0	385	EWLnia amylovera
10	718.5	34.6	338	Hyperresistive res
			358	Hyperresistive res

12	718.5	34.6	338	AAW2407
13	718.5	34.6	338	AAW2407
14	718.5	34.6	338	AAW2407
15	718.5	34.6	338	AAW2407
16	718.5	34.6	338	AAW2407
17	718.5	34.6	338	AAW2407
18	718.5	34.6	338	AAW2407
19	718.5	34.6	338	AAW2407
20	718.5	34.6	338	AAW2407
21	718.5	34.6	338	AAW2407
22	718.5	34.6	338	AAW2407
23	718.5	34.6	338	AAW2407
24	718.5	34.6	338	AAW2407
25	718.5	34.6	338	AAW2407
26	718.5	34.6	338	AAW2407
27	718.5	34.6	338	AAW2407
28	718.5	34.6	338	AAW2407
29	718.5	34.6	338	AAW2407
30	718.5	34.6	338	AAW2407
31	718.5	34.6	338	AAW2407
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34	718.5	34.6	338	AAW2407
35	718.5	34.6	338	AAW2407
36	718.5	34.6	338	AAW2407
37	718.5	34.6	338	AAW2407
38	718.5	34.6	338	AAW2407
39	718.5	34.6	338	AAW2407
40	718.5	34.6	338	AAW2407
41	718.5	34.6	338	AAW2407
42	718.5	34.6	338	AAW2407
43	718.5	34.6	338	AAW2407
44	718.5	34.6	338	AAW2407
45	718.5	34.6	338	AAW2407

## ALIGNMENTS

RESULT 1	AAW563 standard: Protein: 407 AA.
1	AAW563
2	AAW563
3	AAW563
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38	AAW563
39	AAW563
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41	AAW563
42	AAW563
43	AAW563
44	AAW563
45	AAW563

Dislosure; Page 9-10; 75pp; English.

CC This is the amino acid sequence of a 39 kDa, heat stable  
CC hypersensitive response elicitor (HRE) of *Erythra amylovora*. The  
CC control insect on plants are plants are not infected with HRE.  
CC Also claimed is a method of insect control for plants that involves:  
CC (a) providing a transgenic plant or seed transformed with a HRE  
CC (see, e.g., U.S. Pat. No. 5,811,000) and (b) growing the transgenic plants or transgenic  
CC plants produced from the transgenic seeds to control insects. HRE  
CC is a protein that is secreted from the transgenic plants and is  
CC on such plants. It also prevents insects from colonizing host  
CC plants and releasing phytochemicals which result in disease damage to  
CC plants.

Sequence 403 M:

Query Match: 100.0% Score 2079, RA 10, Length 403;  
Best Local Similarity: 100.0% Proct No. 1, 2e155

Matches 403: Conservative 0; Mismatches 0; Indels 0; Gaps 0;

0Y 1 MSATFGLASTWISLIGDAGNNMGLTSSNOMKLGDSNMLGDSNMGDQNTWQGLAEL 60  
0B 1 mslatfglastwislgsdagnmgltsnmgkldgsnmlgdsnmglgdnntwqglael 60  
0B 61 TGNMNMNMWQDGLAGDGLGDSNMLGDSNMLGDSNMLGDSNMLGDSNMLGDSNML 120  
0B 61 tgnnmnmwqdgldagdgldgsnmlgdsnmlgdsnmlgdsnmlgdsnmlgdsnml 120  
0B 61 tgnnmnmwqdgldagdgldgsnmlgdsnmlgdsnmlgdsnmlgdsnmlgdsnml 120  
0Y 121 TTSSTNSPLDGLNFTSTNDSSTSTSTSTSTSTSTSTSTSTSTSTSTSTSTSTST 180  
0B 121 tttstnspldglntstnsdstststststststststststststststststst 180  
0Y 181 GDSGSGDQDPTGDSNMLGDSNMLGDSNMLGDSNMLGDSNMLGDSNMLGDSNML 240  
0B 181 gdsdgsgdqdpdgdsnmlgdsnmlgdsnmlgdsnmlgdsnmlgdsnmlgdsnml 240  
0Y 241 GDSGLDGLSPDYDGLNFTSTNDSSTSTSTSTSTSTSTSTSTSTSTSTSTSTSTST 300  
0B 241 gdsldglspdydglntstnsdststststststststststststststststst 300  
0B 241 gdsldglspdydglntstnsdststststststststststststststststst 300  
0Y 301 GDSGSGDQDPTGDSNMLGDSNMLGDSNMLGDSNMLGDSNMLGDSNMLGDSNML 360  
0B 301 gdsdgsgdqdpdgdsnmlgdsnmlgdsnmlgdsnmlgdsnmlgdsnmlgdsnml 360  
0Y 361 PMAGTGTNGLNMGAGSGLGDSNMLGDSNMLGDSNMLGDSNMLGDSNMLGDSNML 403  
0B 361 pmagtgtnlgmgagsglgdsnmlgdsnmlgdsnmlgdsnmlgdsnmlgdsnml 403

RESULT 2

AMW5455 standard; Protein: 403 AA.

AMW5455:

09-NOV-1998 (first entry)

CC Erythra amylovora hypersensitive response elicitor (HRE).

CC Hypersensitive response elicitor; HRE; growth; transgenic plant.

CC Erythra amylovora.

CC W09832844-AL.

PD 30-JUL-1998.

PD 27-JAN-1998; 98MO-0501507.

27-JAN-1997; 9705-0306048.  
(CORR.) CORNELL RES FOUND INC.

Best SV: Qiu D, Wei Z;

WP01: 1999-427340/36.

N-PDB01: AMW53977.

Method for enhancing plant growth - comprises use of hypersensitive  
P7 response elicitor polypeptide or protein which may also effect, e.g.,  
P8 increase in plant height or earlier germination seed

Dislosure; page 15-16; 10pp; English.

CC This is the deduced amino acid sequence of the 39 kDa, heat stable  
CC hypersensitive response elicitor (HRE) of *Erythra amylovora*. A  
CC method of enhancing growth in plant (compared to) applying a plant  
CC seed under conditions effective to enhance growth of the plant or  
CC plants grown from the seed, or (b) providing a transgenic plant or  
CC polypeptide or protein, and growing the transgenic plant or a plant  
CC produced from the transgenic seed under conditions effective to  
CC enhancing them (see AMW53972-75) can be used to increase plant growth.  
CC The HREs may also result in increased plant height and yield, and  
CC coloration of fruit and plants. E. amylovora HRE can be applied  
CC to tomato plants to enhance growth without causing disease in that  
CC of tomato.

Sequence 403 M:

Query Match: 100.0% Score 2079, RA 10, Length 403;  
Best Local Similarity: 100.0% Proct No. 1, 2e155

Matches 403: Conservative 0; Mismatches 0; Indels 0; Gaps 0;

0Y 1 MSATFGLASTWISLIGDAGNNMGLTSSNOMKLGDSNMLGDSNMGDQNTWQGLAEL 60  
0B 1 mslatfglastwislgsdagnmgltsnmgkldgsnmlgdsnmlgdsnmglgdnntwqglael 60  
0B 61 TGNMNMNMWQDGLAGDGLGDSNMLGDSNMLGDSNMLGDSNMLGDSNMLGDSNML 120  
0B 61 tgnnmnmwqdgldagdgldgsnmlgdsnmlgdsnmlgdsnmlgdsnmlgdsnml 120  
0B 61 tgnnmnmwqdgldagdgldgsnmlgdsnmlgdsnmlgdsnmlgdsnmlgdsnml 120  
0Y 121 TTSSTNSPLDGLNFTSTNDSSTSTSTSTSTSTSTSTSTSTSTSTSTSTSTSTST 180  
0B 121 tttstnspldglntstnsdstststststststststststststststststst 180  
0Y 181 GDSGSGDQDPTGDSNMLGDSNMLGDSNMLGDSNMLGDSNMLGDSNMLGDSNML 240  
0B 181 gdsdgsgdqdpdgdsnmlgdsnmlgdsnmlgdsnmlgdsnmlgdsnmlgdsnml 240  
0Y 241 GDSGLDGLSPDYDGLNFTSTNDSSTSTSTSTSTSTSTSTSTSTSTSTSTSTSTST 300  
0B 241 gdsldglspdydglntstnsdststststststststststststststststst 300  
0B 241 gdsldglspdydglntstnsdststststststststststststststststst 300  
0Y 301 GDSGSGDQDPTGDSNMLGDSNMLGDSNMLGDSNMLGDSNMLGDSNMLGDSNML 360  
0B 301 gdsdgsgdqdpdgdsnmlgdsnmlgdsnmlgdsnmlgdsnmlgdsnmlgdsnml 360  
0Y 361 PMAGTGTNGLNMGAGSGLGDSNMLGDSNMLGDSNMLGDSNMLGDSNMLGDSNML 403  
0B 361 pmagtgtnlgmgagsglgdsnmlgdsnmlgdsnmlgdsnmlgdsnmlgdsnml 403

RESULT 3

AMW6114 standard; Protein: 403 AA.

AMW6114:















XX Ewlnia chrysanthemi.

XX Key Location/Qualifiers

XX Misc-difference 19 /note= "encoded by GCGC00907"

XX WO20002905-42.

XX 18-MAY-2000.

XX 04-MAY-1999: 99MO-00360139.

XX 05-MAY-1998: 98MS-0197431.

XX (EHRN-) EDRN BIOSCIENCE CORP.

XX Mel Z, Schaling LJ.

XX WPI: 2000-37556/72.

XX N-PDB: AAO00657.

XX Application of a hypersensitive response elicitor protein to plants to

XX impact stress resistance

XX disclosure: Page 4-5; kapp: English.

XX The patent discloses a method to impart stress resistance to plants by

XX response elicitor protein from Ewlnia chrysanthemi. The protein is meant

XX stable and used to impart stress resistance to plants.

XX Sequence 318 AA:

XX Query Match Similarity 31.48; Score 718.5; DB 21; Length 318;

XX Matches 173; Conserved: 41; Mismatches 111; Indels 79; Gaps 11;

XX 13 MOIS-GAGGNCMLGTSQNKAGC-NSK---LGLGSCNNDNTVQACLTQKQNMW 67

XX 1 mqltlahlgdlgyagvqag-ghlmaasasvavhlaetlclatlam-- 55

XX 68 SMOGCGIAGGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGL 127

XX 56 -----gagllagvqag-ghlmaasasvavhlaetlclatlam-- 96

XX 128 PLDGLGINSQNDSTGSGSGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLG 179

XX 97 -----gagllagvqag-ghlmaasasvavhlaetlclatlam-- 128

XX 180 TQSSGSGGPTFGSDWAKYKGTALGSLGMLGSLGGLGGLGGLGGLGGLGGLGGLG 239

XX 129 laahlaahagmhlqgmhlqgmhlqgmhlqgmhlqgmhlqgmhlqgmhlqgmhlqgm 174

XX 240 LGLGSCNNDNTVQACLTQKQNMW 299

XX 175 19agvqagvqagvqagvqagvqagvqagvqagvqagvqagvqagvqagvqagv 334

XX 300 TQSSGSGGPTFGSDWAKYKGTALGSLGMLGSLGGLGGLGGLGGLGGLGGLGGLG 359

XX 235 19agvqagvqagvqagvqagvqagvqagvqagvqagvqagvqagvqagvqagv 384

XX 360 RMAVQAGINSQNDSTGSGSGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLG 403

XX 295 avagvqagvqagvqagvqagvqagvqagvqagvqagvqagvqagvqagvqagv 358

XX Result 1A

XX ID AAH4853 standard Protein: 318 AA.

XX AAH4853

XX

AC AAH4853;

XX 08-AUG-2000 (first entry)

XX A hypersensitive response elicitor protein.

XX Hypersensitive response; insect control; disease resistance;

XX hypersensitive response elicitor; plant growth; vegetable; crop;

XX ornamental plant.

XX Ewlnia chrysanthemi.

XX Key Location/Qualifiers

XX Misc-difference 20 /note= "encoded by GCGC00907"

XX WO20002905-42.

XX 13-APR-2000.

XX 05-OCT-1998: 99MO-00360139.

XX N-PDB: AAO00657.

XX (EHRN-) EDRN BIOSCIENCE CORP.

XX Mel Z, Fan H, Magermeier JI.

XX WPI: 2000-303745/76.

XX N-PDB: AAO14937.

XX Hypersensitive response elicitor polypeptides useful for imparting

XX enhanced growth, disease resistance and insect resistance to plants,

XX especially vegetables and ornamental flowers.

XX disclosure: Page 6-7; 10pp: English.

XX The present sequence represents a hypersensitive response elicitor

XX polypeptide. The specification describes hypersensitive response

XX elicitor polypeptide fragments, which do not elicit a hypersensitive

XX response in plants, and/or control insects. The polypeptide

XX fragments may be used to these properties to plants. The plants which

XX may be treated in this way include, but are not limited to, tomato,

XX cucumber, eggplant, pepper, zucchini, pumpkin, squash, melon,

XX endive, cabbage, brussels sprout, bean, pea, chori, lettuce,

XX peanut, corn, potato, sweet potato, bean, pea, chori, lettuce,

XX carrot, squash, pumpkin, zucchini, cucumber, apple, pear, melon,

XX citrus, strawberry, grape, raspberry, pineapple, soybean, tobacco,

XX petunia, platagium, poinsettia, chrysanthemum, carnation or stinla.

XX Sequence 318 AA:

XX Query Match Similarity 31.48; Score 718.5; DB 21; Length 318;

XX Matches 173; Conserved: 41; Mismatches 111; Indels 79; Gaps 11;

XX 13 MOIS-GAGGNCMLGTSQNKAGC-NSK---LGLGSCNNDNTVQACLTQKQNMW 67

XX 1 mqltlahlgdlgyagvqag-ghlmaasasvavhlaetlclatlam-- 55

XX 68 SMOGCGIAGGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGL 127

XX 56 -----gagllagvqag-ghlmaasasvavhlaetlclatlam-- 96

XX 128 PLDGLGINSQNDSTGSGSGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLGGLG 179

XX 97 -----gagllagvqag-ghlmaasasvavhlaetlclatlam-- 128

XX 180 TQSSGSGGPTFGSDWAKYKGTALGSLGMLGSLGGLGGLGGLGGLGGLGGLGGLG 239





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Thu May 2 11:45:16 2002

us-09-770-693-3, Raj

Page 2

[illegible][illegible]

































































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OM protein - protein search, using SW model

Run on: May 2, 2002, 03:32:15 / Search Time 39.55 Seconds

196,199 Million Cell updates/sec

File: us-070-693.3  
Sequence: 1 NLSLSQSLASPMQISGIA.....DAWAGDINNALDIA 403

Scoring table:  
BLOSUM62  
Gap: 10.0, Expect: 0.5

Searched: 473505 seqs, 146272329 residues

Total number of hits satisfying chosen parameters: 473505

Minimum DB seq length: 0

Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%  
Maximum Match 100%

Listing first 45 summaries

Database: 1: sp.archaea;  
2: sp.bacteria;  
3: sp.eukarya;  
4: sp.human;  
5: sp.lavatebacte;  
6: sp.mycobact;  
7: sp.micr;  
8: sp.mycobact;  
9: sp.mycobact;  
10: sp.mycobact;  
11: sp.mycobact;  
12: sp.mycobact;  
13: sp.mycobact;  
14: sp.unclassified;

Prod. No. is the number of result predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

# SUMMARIES

Result	Score	Query Length	DB ID	Description
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2	1181.5	56.8	382	01-07-01-2001 (TRENDEL, 16, Created)
3	1181.5	56.8	382	01-07-01-2001 (TRENDEL, 16, Created)
4	717.2	34.2	142	01-07-01-2001 (TRENDEL, 16, Created)
5	717.2	34.2	142	01-07-01-2001 (TRENDEL, 16, Created)
6	245.5	12.3	591	007224 mycobact
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18	245.5	12.3	591	007224 mycobact
19	245.5	12.3	591	007224 mycobact

20	219.5	11.0	837	2	051684	051684 mycobact
21	229.5	11.0	1329	2	006810	006810 mycobact
22	229.5	11.0	486	10	09A823	09A823 mycobact
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## ALIGNMENTS

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[illegible][illegible]



























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 ACCESSION F001897.1 GI:397974  
 KEYWORDS  
 SOURCE Unknown  
 ORGANISM Unclassified.  
 REFERENCE 1 (bases 1 to 1158)  
 TITLE Hyperimmune response induced resistance in plants  
 JOURNAL Patent: US 5776889-A 4 07-JUL-1998.  
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 ORIGIN

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FH Key Location/Qualifiers  
FT 71..1284  
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X3  
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X6 03-DEC-1998.  
X7  
X8 28-MAY-1998: 9800-0510874.  
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X10 30-MAY-1997: 9705-0048199.  
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X12 (SDS) : DEBN BIOSCIENCE CORP.  
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X15  
X16 WPI: 1999-070210/05.  
X17 P-250: M6861693.  
X18  
X19 New fragments of an Ewalia hypersensitive response elicitor protein  
X20 and related DNA used to impart disease resistance to plants, to  
X21 increase their growth and to control insects  
X22  
X23 Disclaimers: Page 12: 9dpp: English.  
X24  
X25 The present sequence encodes a hypersensitive response elicitor  
X26 protein (also called harpin protein) that is able to elicit a  
X27 hypersensitive response in plants. The protein is able to induce  
X28 hypersensitive response elicitors from other pathogenic organisms.  
X29 The protein, in non-infectious form, is applied to plants to impart  
X30 disease resistance to a wide range of plants.  
X31 The protein is also used to increase the quantity and quality of seeds,  
X32 to provide earlier germination etc.) and to control insects (e.g. corn  
X33 borers, Lepidoptera larvae etc.) The same results are provided by  
X34 transgenic plants expressing the protein.  
X35  
X36 Sequence 1288 BP: 309 A: 297 C: 128 G: 254 T: 0 other:  
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X38  
X39  
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Hyper-sensitive response elicitor; environmental stress resistance;  
 plant; 2s.  
 Evidna amyDvora.  
 Reg Location/Qualifiers  
 CDS /feature  
 /product="Hyper-sensitive response elicitor"  
 MO00028055:42.  
 XX 18-MAY-2000.  
 PE 04-NOV-1993; 9900-US26039.  
 PA 05-NOV-1998; 9805-0107343.  
 XX (EENR) EENR BIOLOGICAL COMP.  
 PA Malt Z, Solating BL.  
 XX WPI: P500-375556/32.  
 DX P-RESD: AAV71093.  
 DX Application of a hyper-sensitive response elicitor protein to plants to  
 XX elicit stress resistance.  
 XX The present disclosure a method to impart stress resistance to plants by  
 XX applying a hyper-sensitive response elicitor in a non-infectious form to  
 XX response elicitor protein construct. The construct is a DNA encoding hyper-sensitive  
 XX The present sequence is used to transform transgenic plant or plant  
 XX seeds to impart stress resistance.  
 XX Sequence 1288 BP; 309 A; 297 C; 428 G; 254 T; 0 other:  
 Query Match 100.0%; Score 1288; DB 21; Length 1288;  
 Best Local Similarity 100.0%; Pval No. 0;  
 Matches 1288; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

DB 421 gggggaacatccctcccaacaacatcccgccgacacggcgccggggtctaac 480  
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 QY 481 tgaagatcccaaaagagagagagagagagagagagagagagagagagagag 540  
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 DB 481 tcaagctcccaaaagagagagagagagagagagagagagagagagagagag 500  
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 QY 541 cccgagtcgagctgagagagagagagagagagagagagagagagagagag 600  
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 DB 541 ccagatcgagctgagagagagagagagagagagagagagagagagagag 560  
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 QY 601 caagatcgagctgagagagagagagagagagagagagagagagagagag 660  
 |||||||  
 DB 601 caagatcgagctgagagagagagagagagagagagagagagagagagag 660  
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 QY 661 gctccatcaaaaagagagagagagagagagagagagagagagagagagag 720  
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 DB 721 cctccatcgag 780  
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 QY 781 gctccatcgag 840  
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 DB 781 gctccatcgag 840  
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 QY 841 ttagatgag 900  
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 DB 901 atcgatcaacgag 960  
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 QY 961 gcaag 1020  
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 QY 1021 cgaag 1080  
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 QY 1141 atgatatcaag 1200  
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 QY 1201 ggtggtctcgag 1260  
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 DB 1201 ggtggtctcgag 1260  
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 QY 1261 ctggcag 1288  
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 DB 1261 ctggcag 1288  
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RESULT 6  
 ID AAL4938 standard; DNA; 1288 BP.  
 XX AAL4938:  
 XX 08-MAY-2000 (files entry)  
 PE DNA encoding a hyper-sensitive response elicitor protein.  
 DE Hyper-sensitive response elicitor control; disease resistance;  
 KM Hyper-sensitive response elicitor; plant growth; vegetable crop;  
 ornamental plant; 2s.

















































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1 TITLE OF INVENTION: Elicitor of the hypersensitive response in plants
2 NUMBER OF SEQUENCES: 5
3 COMPLETER: Abasco, associates
4 ADDRESS: Abasco, associates
5 STREET: 25 Ayrton Drive
6 CITY: Tulum, Quintana Roo
7 STATE: Quintana Roo
8 COUNTRY: USA
9 ZIP: 06611
10 COMPLETER PHONE:
11 MEDIUM TYPE: floppy disk
12 SOFTWARE: Microsoft DOS
13 SOFTWARE: Microsoft DOS
14 CURRENT APPLICATION DATA: 4.0
15 APPLICATION NUMBER: PCT/US93/06243
16 CLASSIFICATION: 5930630
17 PRIORITY DATA:
18 PRIORITY DATE: 01-11-1992
19 FILING DATE: 01-11-1992
20 ATTORNEY/AGENT INFORMATION:
21 NAME: George M. Knaack
22 TELEPHONE: 874
23 TELEFAX: 874
24 REFERENCE/DOCKET NUMBER: CWR-D-1172
25 TELEPHONE: 874
26 INFORMATION FOR SEQ ID NO: 1:
27 LENGTH: 1158 base pairs
28 TYPE: nucleic acid
29 TOPOLOGY: linear
30 MOLECULE TYPE: DNA
31 PCT-0593-06243-4
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